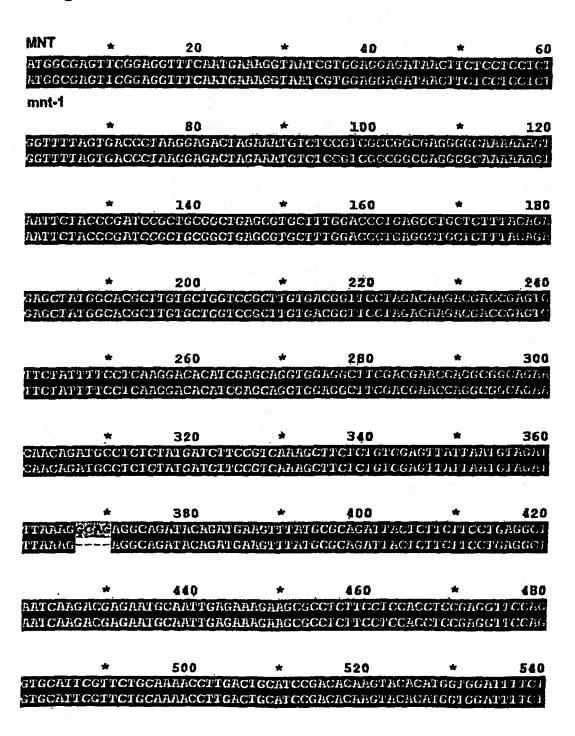
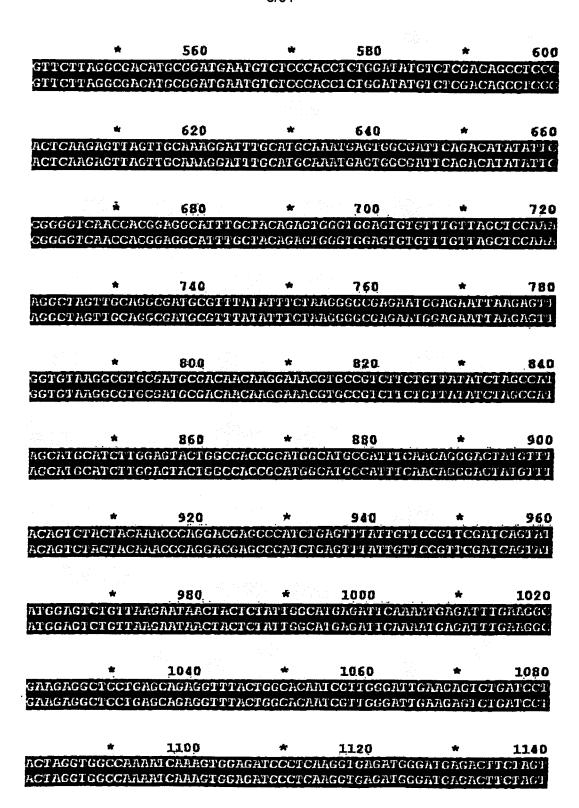
REPLACEMENT SHEET 7/51

Figure 6

Alignment of w.t. MNT and mutant mnt-1 cDNA



REPLACEMENT SHEET 8/51



REPLACEMENT SHEET 9/51

	*	1,160	**	1180	*	1200
		ATAGAGTATCT				
ATTCCT	CGACCTG	ATAGAGTATCT	CCGTGGNAA	GTAGAGCCAG	CHCTHGCTCC	Techeci
	*	1220	*	1240	*	1260
		CAALGCCTAGG				
TTGAGT	CCTGTTC	CHATGCCTAGG	CCTARGAGG	ecchent <i>en</i> a	AHATAGCACC	THEATCH
	*	1280	*	1300	*	1320
		TGCTTACCAGA				
CCIGRO	ICTICGA	TGCTTACCAGA	GAAGGTACA	ACTVITAGGOIVA	ACATECHECO	A MATICOL
	*	1340	*	1360	*	1380
		CARGGGTCTTG				
GCMIGC	CHCHI	CHAGGGTCTTG	CALEGICAL	CHAIN VACITORY	COMMERCE	GRUMAURU
	. *	1400	*		*	1440
		AGTGTGATGCT				
ACTGAGE	GI GIVE	AGTGTGATGCT	SOLEWRING TANK	TOTGITTETOT	GGUAAT CITTU	たほしけけれ
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		AACCTACTTAC				
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		gtgattcagaa				
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		PCTCCCTGAAG				

REPLACEMENT SHEET 10/51

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	*	1940	*	1960	*	1980
		AGACAGCAAA				
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	*	2000	ja je 🔅 😼	2020	*	2040
		TGAATGGGACA				
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		ichendra 2.5c.				
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		2120	# CGTGRACAG	GGAAGACCAT	TCCAGACTAA	2160 Tantcol
		2120 CAAACGATCAT CAAACGATCAT	# CGTGRACAG	GGAAGACCAT GGAAGACCAT	TCCAGACTAA	2160 PANGCI PAN CC1
GGT CA	ARRTCHA	2120 Charcgateat Charcgateat 2180	* CGTGRACAG CGTGRACAG	GGAAGACCAT GGAAGACCAT 2200	TCCAGACTAA TCCAGACTAA	2160 TWATEGI TWATEGI 2220
GGTCA	ARRTCAN * Anggatg	2120 CAAACGATCAT CAAACGATCAT	* ICGTGRACAG ICGTGRACAG * IACCAACTCA	GGAAGACCAT GGAAGACCAT 2200 BGTAGGAGTT	TCCAGACTAA TCCAGACTAA * GCACAAAGGT	2160 TANGCI TANI CCI 2220 TONGNA
GGGTCA CATCCG	ARRTCAN * Anggatg	2120 CAMACGATCAT CHARCGATCAT 2180 CTCMBACGAM	* ICGTGRACAG ICGTGRACAG * IACCAACTCA	GGAAGACCAT GGAAGACCAT 2200 BGTAGGAGTT	TCCAGACTAA TCCAGACTAA * GCACAAAGGT	2160 TANGCI TANI CCI 2220 TONGNA
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GGTCA CATCCG	ARATICAA * AAGGATG AAGGATG	2120 CAMACGATCAT CHARCGATCAT 2180 CTCMBACGAM	CGTGRACAG CGTGRACAG * ACCRACTCR ARCCRACTCR	GGAAGACCAT GGAAGACCAT 2200 AGTAGGAGTT AGTAGGAGTT	TCCAGACTAA TCCAGACTAA TCCAGACTAA GCACAAAGGT GCACAAAGGT	2160 TMATCCI TMATCCI 2220 TCACAAG TCACAAG
GGTCA CATCCG CATCCG	ARRICAA * AAGGATG AAGGATG * ATTGCAC	2120 CAAACGATCAY CHAACGATCAY 2180 CTCAAACGAAY CTCAAACGAAY	* ICGTGRACAG ICGTGRACAG * ACCAACTCA ARCGARCTCA	GGAAGACCAT GGAAGACCAT 2200 BGTAGGAGTT AGTAGGAGTT 2260 TCAAAGTICC	TCCAGACTAA TCCAGACTAA TCCAGACTAA CCACAAAGGT GCACAAAGGT	2160 TWATCCI TWATCCI 2220 TCACARG TCACARG
GGTCA CATCCG CATCCG	ARRICAA * AAGGATG AAGGATG * ATTGCAC	2120 CARACGATCAT CHARCGATCAT 2180 CTCARACGARA CTCARACGARA 2240 TTGGCCGTTCA	* ICGTGRACAG ICGTGRACAG * ACCAACTCA ARCGARCTCA	GGAAGACCAT GGAAGACCAT 2200 BGTAGGAGTT AGTAGGAGTT 2260 TCAAAGTICC	TCCAGACTAA TCCAGACTAA TCCAGACTAA CCACAAAGGT GCACAAAGGT	2160 TWATCCI TWATCCI 2220 TCACARG TCACARG
GGTCA CATCCG CATCCG	ARRICAA * AAGGATG AAGGATG * ATTGCAC	2120 CARACGATCAT CHARCGATCAT 2180 CTCARACGARA CTCARACGARA 2240 TTGGCCGTTCA	* ICGTGRACAG ICGTGRACAG * ACCAACTCA ARCGARCTCA	GGAAGACCAT GGAAGACCAT 2200 BGTAGGAGTT AGTAGGAGTT 2260 TCAAAGTICC	TCCAGACTAA TCCAGACTAA TCCAGACTAA CCACAAAGGT GCACAAAGGT	2160 TWATCCI TWATCCI 2220 TCACARG TCACARG

REPLACEMENT SHEET 11/51

	*	2360	*	2380	*	- 2400
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					GTTGGTGACGR	
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CAGGAG!	TTTTGTT	GCATGGTTCG	CRAAATCTTC	ATATACACG	ARAGREGARGT	GAGGAAG
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					rgcgctgggar rgcgctgggra	
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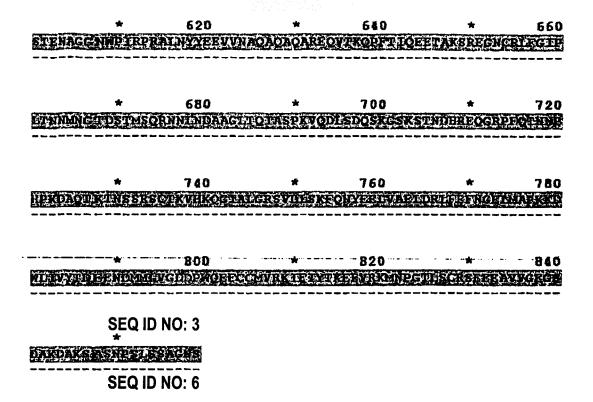
REPLACEMENT SHEET 12/51

Figure 7

Alignment of w.t. MNT and mutant mnt-1 protein

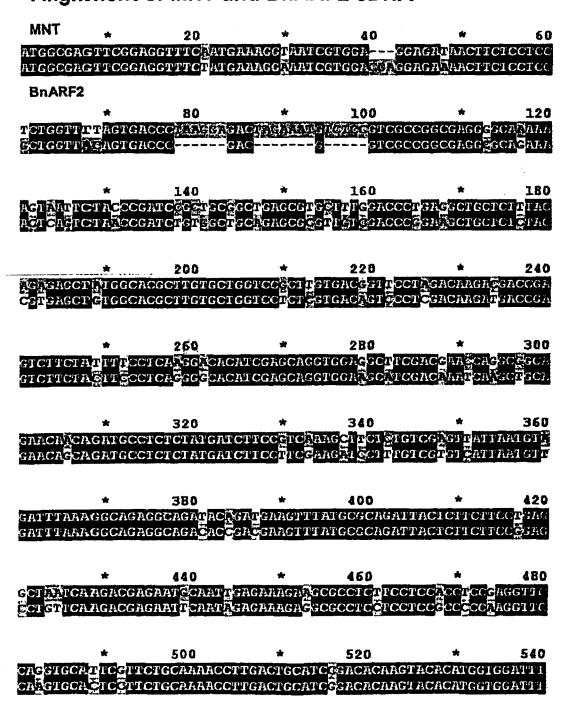
MNT	*	20	•	40	•	60
MASSEY	SMEGHRG	GDNFSSSGFSD	DKETEINSVI		SALAFDALDI	
MASSEY	SMKGNRG	GDNFSSSGFSD	PKETRIVSVI	GEGOKSUSTR	SABAFRALDI	IE ADT VE
mnt-1						
1,1116						
	*	80	*	100	*	120
		PRODDRYFYFD				
ELWHAC	AGPLVIVI	PRODDRVFYFP	OCHIEQVENS	THOAREQUE	LYDLPSKLLC	EVINVD
				•		
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	*	200	*	220	*	240
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BESMI	HEE WOKA	NMOP RPASETS	RVIOCORYS	FRIKATERVI	COMPENSIV	OSSAI
	*	500	*	520	*	540
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	*	560	*	580	*	600
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REPLACEMENT SHEET 13/51



REPLACEMENT SHEET 14/51

Figure 8 Alignment of MNT and BnARF2 cDNA



REPLACEMENT SHEET 15/51

	*	560	*	580	*	600
TOTETT CT	TAGGCGA	CatgCggatgai CatgCggatgai	ATGTCTCCC	ACCTCTGGATA	TGTCTCGAC	AGCCI
i et et éct	TAGGCG S	CATGCGGATGAT	MGTCTCCC	ACCICIEGAIA	TO TO YOUR	Witter
	*	620	*	640	*	660
CCGACTCA	GAGTTA	GTTGC/AAAGGAV	FIGCATGO	NATTARGEGE	GATTICAGAC	MAKER
CCHRCTCR	General V	стт <u>ссала Я</u> сат	FETGCATGC	nn <mark>g</mark> gretege	GRITCEGAC	TATA
		cna.		700		701
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		CenageCattic				
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ar <mark>c</mark> agect	GETGECA	GGCGATGCTTT	ATATTTCT	AAGGGGGGAGAI	ATGGAGAAT	TIP COL
		ma.a	بغ	DOA		D 4 6
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st <mark>e</mark> sstet	AAGGCGT	GCZNTGCGGCNG	CRAGGAAA	received in	CIGTTATAT	e Jage
	•	860	*	880	•	900
CATAGGAI	GCATCT I	GGRGTA SIGGCO			CANCACCE	
		CONCLATINGCO				
	•	920	*	940	*	960
TTTACAGT	CTACTA	NANCC BAGGACE				
TTTACAGI	CI ACI AT	ARACC <mark>GAGGRO</mark> R	REPERTIC	A GAGTTTATTGI	TECGITTE	ai cag
	*	980	*	1000	•	1020
TATATGGA	GICTSPT	RAGARINACTAC	ICTATUSG	Catgagatt e al	iamgagan	TTGAR
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	*	1040	*	1060	*	1080
		GAGCAGAGGTTT				TGAT
GGCGAAGA	GCTCC.	CACCAGAGGTTT	ACTGGCAC	AATCGTTGGGAT	Теллеліўн	CTGA
	*	1100	*	1120	*	1140
CCTACTAG	STGG STA	AAATCAAAGIGG	AGATCCCTC	CHAGGIGAGAIG	GGATGAGAG	TriOI

REPLACEMENT SHEET 16/51

	*	1160	*	1180	*	1200
ACTATT	CCTCGA	CTGATAGAGT	ATCTCCGTGG	AHAGIIAGAGCC	ACCICITEC	TCCTCCT
AGTATTO	CCTCC	CCTGATAGAGT	ate <mark>é</mark> ccetes	AR <mark>GR</mark> TAGAGCC	AGCTCTTTC	TCCTCCT
	<u>*</u>	1220	*	1240	*	1260
				AGGCCCAGATC		
GCTTTG	AGGCCTC	TRECHATECC	TAGGCCTAAG	AGGCCCAGATC	TARTCIAGO	TT CIVICA
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	*	1340	*	1360	*	1380
CCACCII	GGGAC		CHARGE SERVICE COLUMN	CHAGNAVACTE	GACCITICAC	
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	*	1520	*	1540	*	1560
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		ATTGCACTUG				
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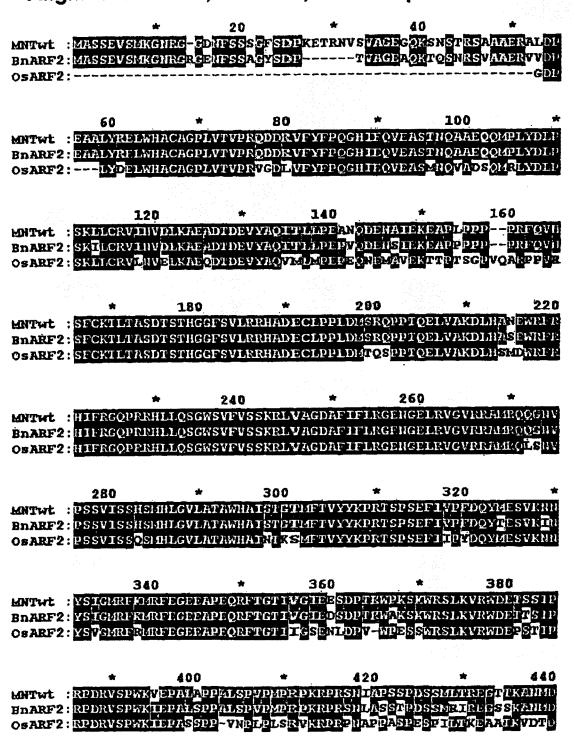
REPLACEMENT SHEET 18/51



SEQ ID NO: 55
ACTOTPAA
ACTOTPAA
SEQ ID NO: 9

REPLACEMENT SHEET 19/51

Figure 9 Alignment of MNT, BnARF2, OsARF2 proteins



REPLACEMENT SHEET 20/51

